

**Metabolic Engineering of  
*Escherichia coli* W3110 for  
Redox Neutral & Oxidized Products**

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# Agricultural Opportunity - BioRefinery



## Biomass Feedstock

- Trees
- Grasses
- Agricultural Crops
- Agricultural Residues
- Animal Wastes
- Municipal Solid Waste

## Conversion Processes

- Enzymatic Fermentation
- Gas/liquid Fermentation
- Acid Hydrolysis/Fermentation
- Gasification
- Combustion
- Co-firing

## USES

### Renewable Fuels:

- Ethanol
- Bio-Diesel

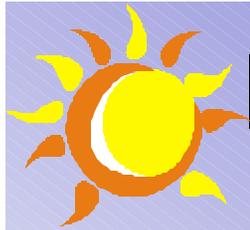
### Renewable Power:

- Electricity
- Heat or CHP

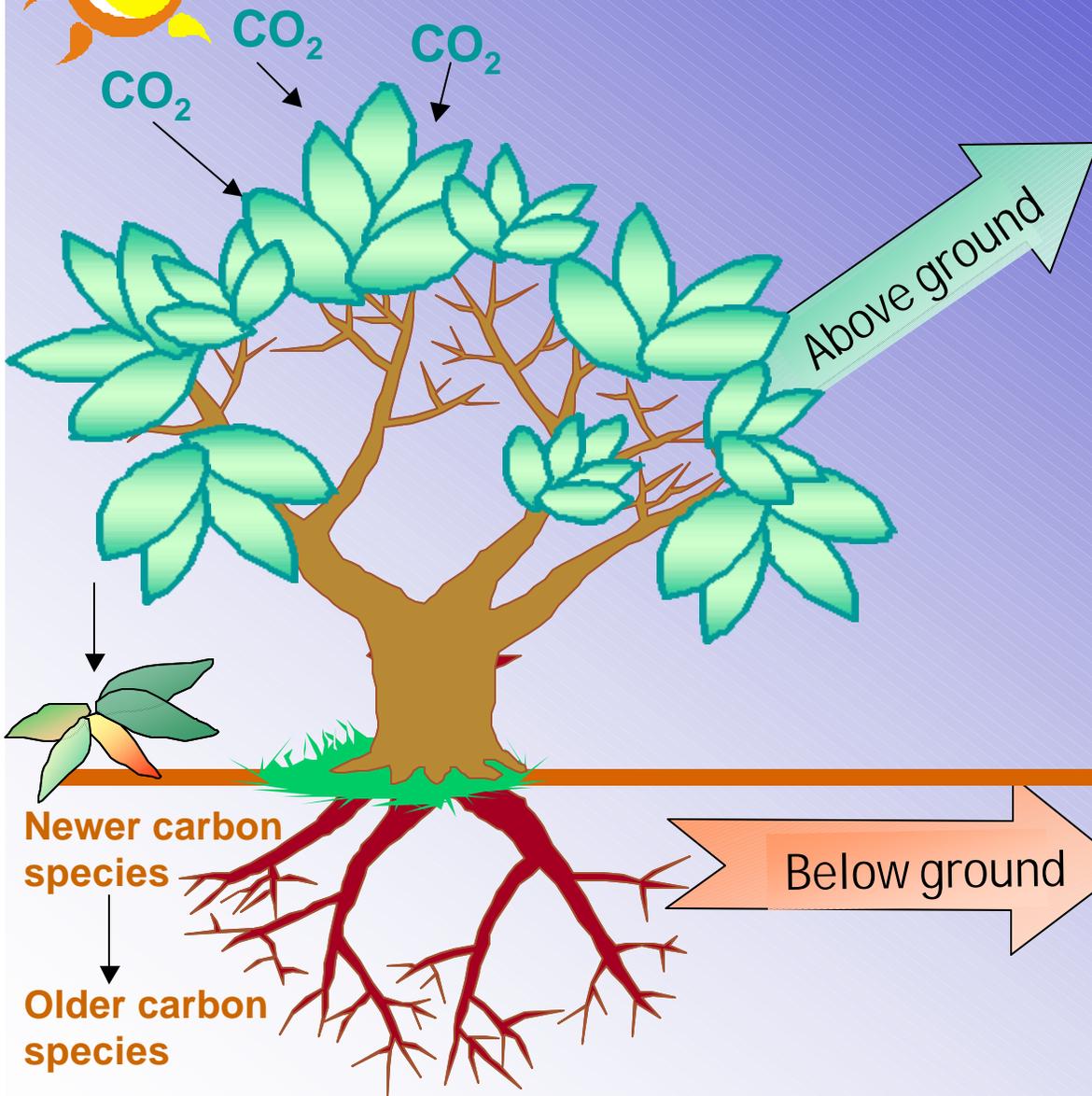
### Renewable Chemicals

- Plastics
- Solvents
- Chemical Intermediates
- Phenolics
- Adhesives
- Furfural
- Fatty acids
- Acetic Acid
- Carbon black
- Paints
- Dyes, Pigments, and Ink
- Detergents
- Etc.

### Food and Feed and Fiber



# Renewable Fuels and Chemicals:



## ❖ Displacement of oil

- **Commodity chemicals**
  - polylactic acid
  - 3-HP, 1-3 PD
  - solvents
  - acids
- **Fuels**
  - ethanol
  - biodiesel
  - power
- **Rural Employment**

## ❖ **Carbon Sequestration of in soil (short term)**

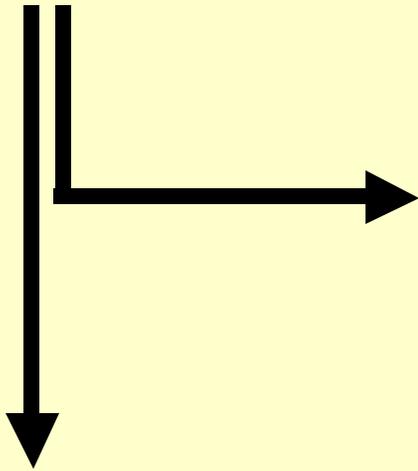
# **What are the limitation of Metabolic Engineering in *E. coli*?**

- 1. Can we combine the beneficial features of aerobic and anaerobic metabolism?**
  - **Aerobic:** High growth rate, low NADH, external electron acceptor ( $O_2$ )
  - **Anaerobic:** High glycolytic flux, low  $CO_2$  production, low cell yield, high product yield
- 2. As an example, glucose to acetate (pyruvate)**

# Overview of Metabolism

**Aerobic, + oxygen**

**Glucose,  $C_6H_{12}O_6$**

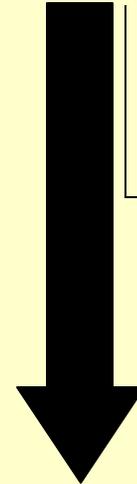


**Cell Mass  
50% of Carbon**

**$CO_2 + 36 \text{ ATP (max)}$   
50% of carbon**

**Anaerobic, - oxygen**

**Glucose,  $C_6 H_{12} O_6$**



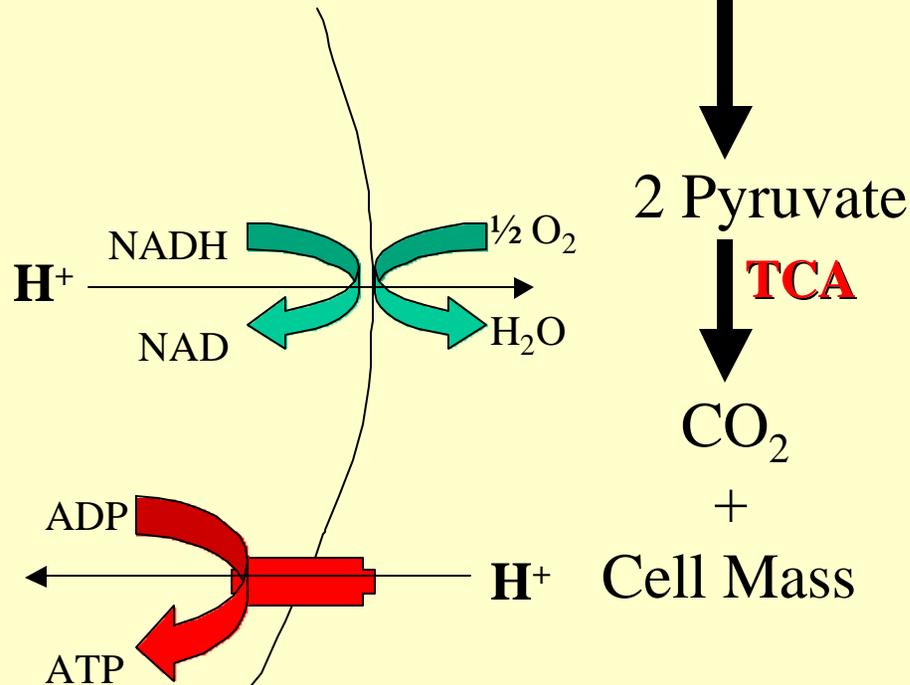
**Cell Mass  
5% of Carbon**

**Fermentation Products + 2 ATP  
95% of Carbon  
(most commodity products)**

# Overview of the Problem

## Aerobic, + oxygen

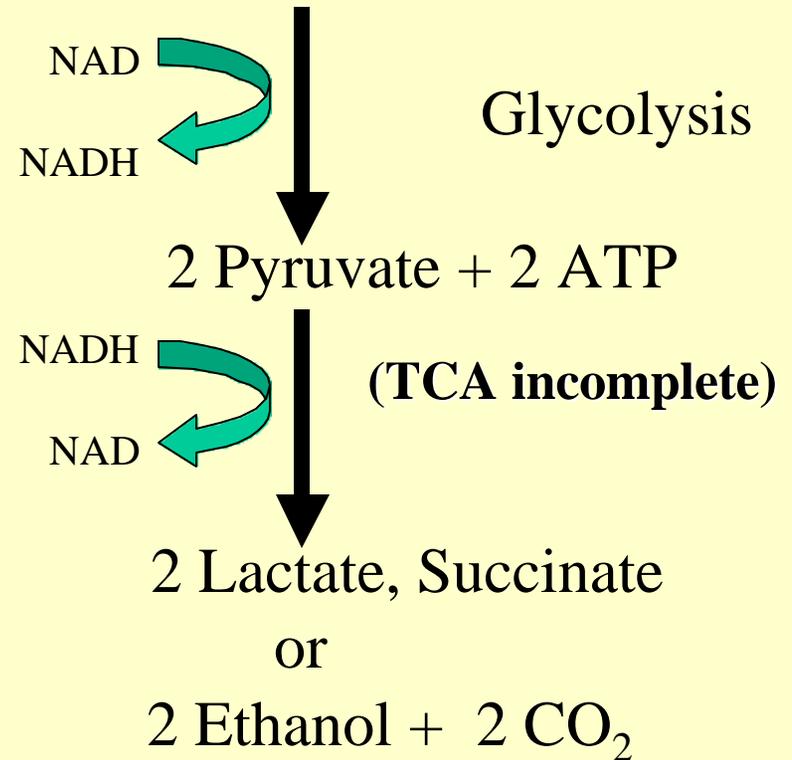
Glucose,  $C_6H_{12}O_6$



$CO_2 + 36$  ATP (max)  
50% of carbon as cell mass  
50% of carbon as  $CO_2$

## Anaerobic, - oxygen

Glucose,  $C_6H_{12}O_6$



Fermentation Products + 2 ATP  
5% of carbon as cell mass  
95% of carbon as product

# Three Problem Areas

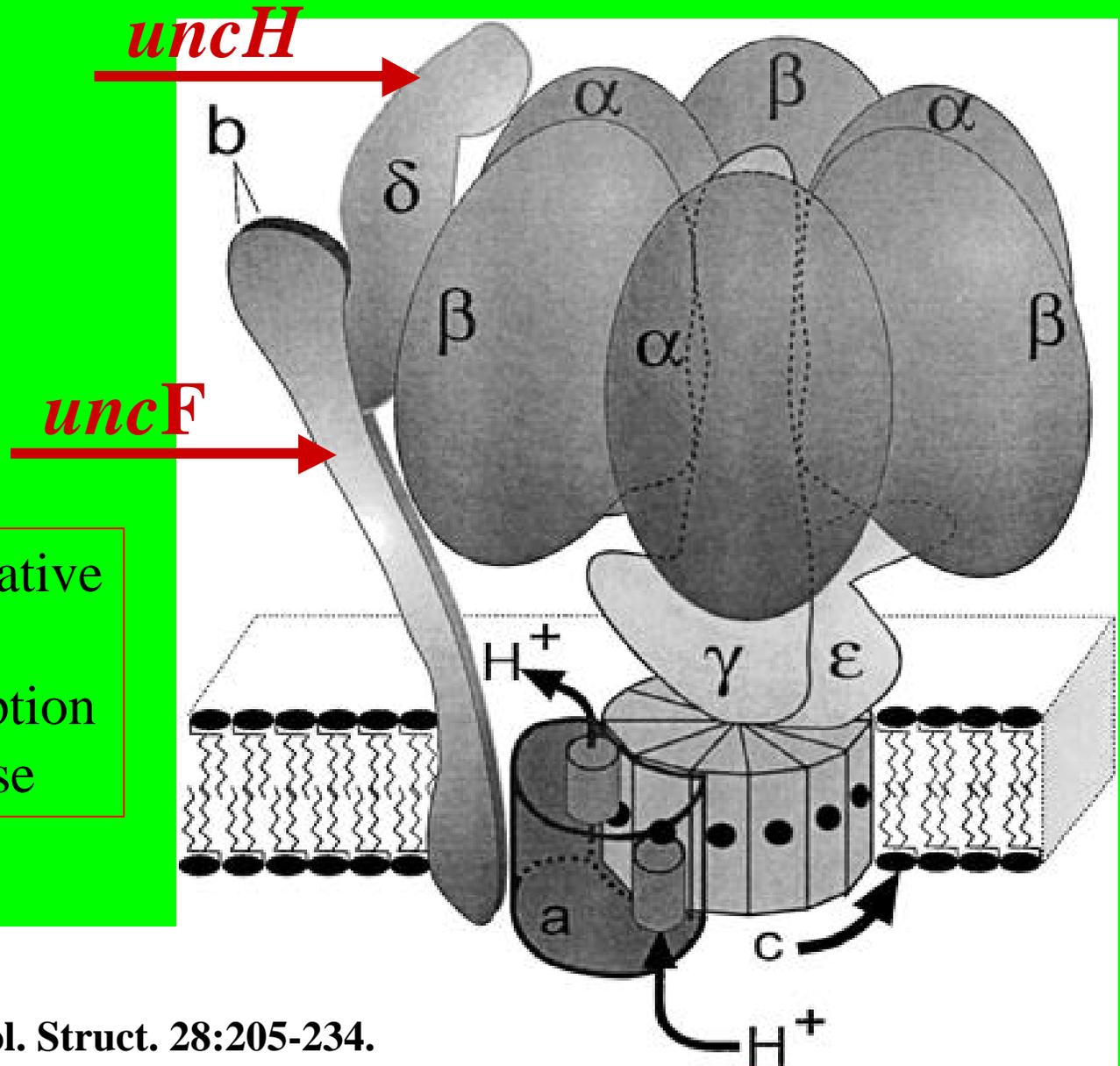
1. Cells → Too many – reduce ATP
2. Volatiles → Too much carbon lost  
– interrupt TCA cycle, inactivate ADH
3. Fermentation products → potential sink  
– inactivate pathways

“Just  
turn  
me  
loose!”

Oxidative Phos negative  
Gratuitous consumption  
of ATP by F<sub>1</sub>ATPase

Nakamoto et al., 1999

Ann. Rev Biophys. Biomol. Struct. 28:205-234.



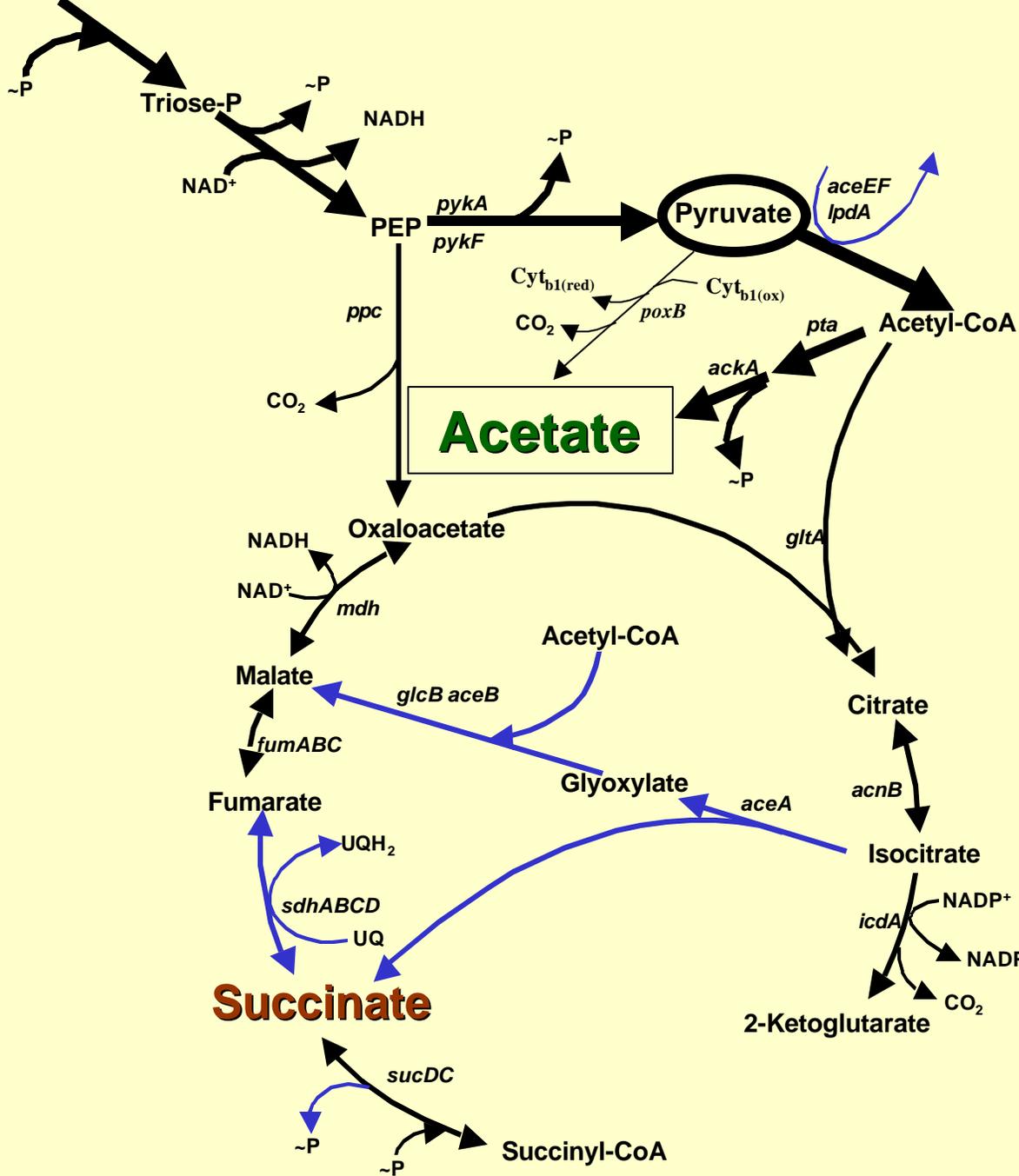


# Strain Construction

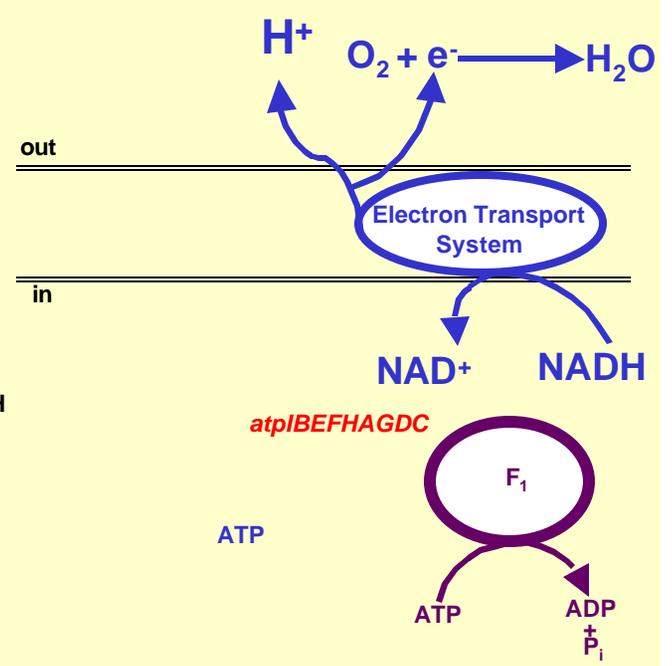
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1. Homologous recombination, confirmed by PCR and sequence, one mutation per strain
2. Phage P1 to combine mutations into single strain
3. Flanking FRT sites and FLP recombinase to remove antibiotic markers used for selection
4. Fusaric acid selection to create deletions from Tn10 insertions
5. PCR cloning and sequencing to confirm each step

1/2 Glucose



—> Both  
—> Aerobic only  
—> Anaerobic only



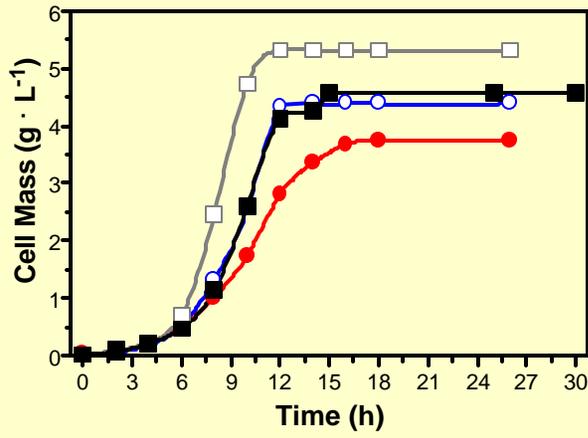
*atpIBEFHAGDC*

ATP

F<sub>1</sub>  
ATP  
ADP + P<sub>i</sub>

# Fermentation Conditions:

- mineral salts, 37°C
  - 3 % glucose
  - starting  $OD_{550} = 0.1$
  - 1% inoculum
  - 10 L initial volume
  - agitation set @ 450 rpm
  - pH controlled @ 7.00 with 45% w/w KOH (11.4 M)
  - DO controlled @ 5% of air saturation by adjusting the ratio of air, O<sub>2</sub> and N<sub>2</sub>
- Gas flow 1 L/min (0.1vvm)

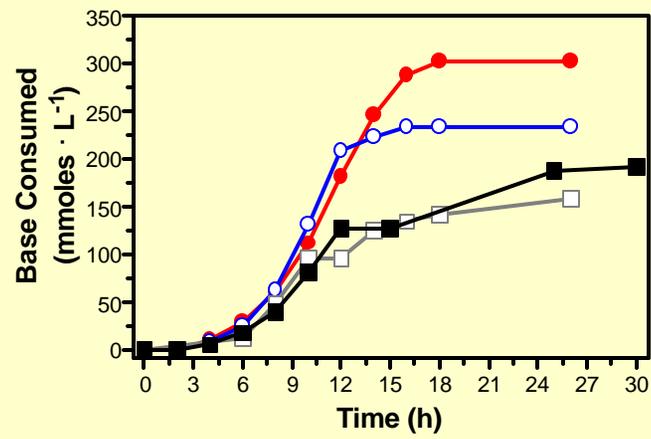
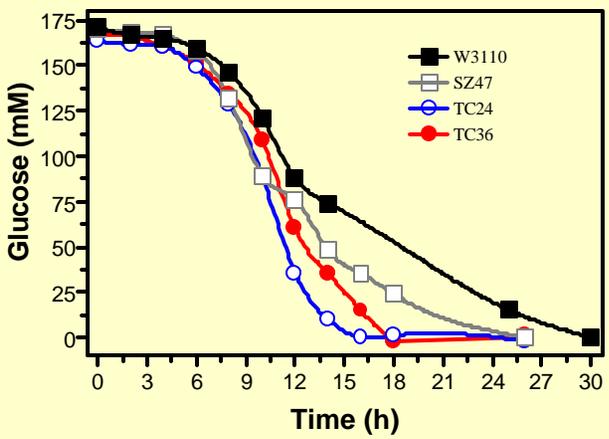


Strain W3110 (wild type) ~30 ATP/glu

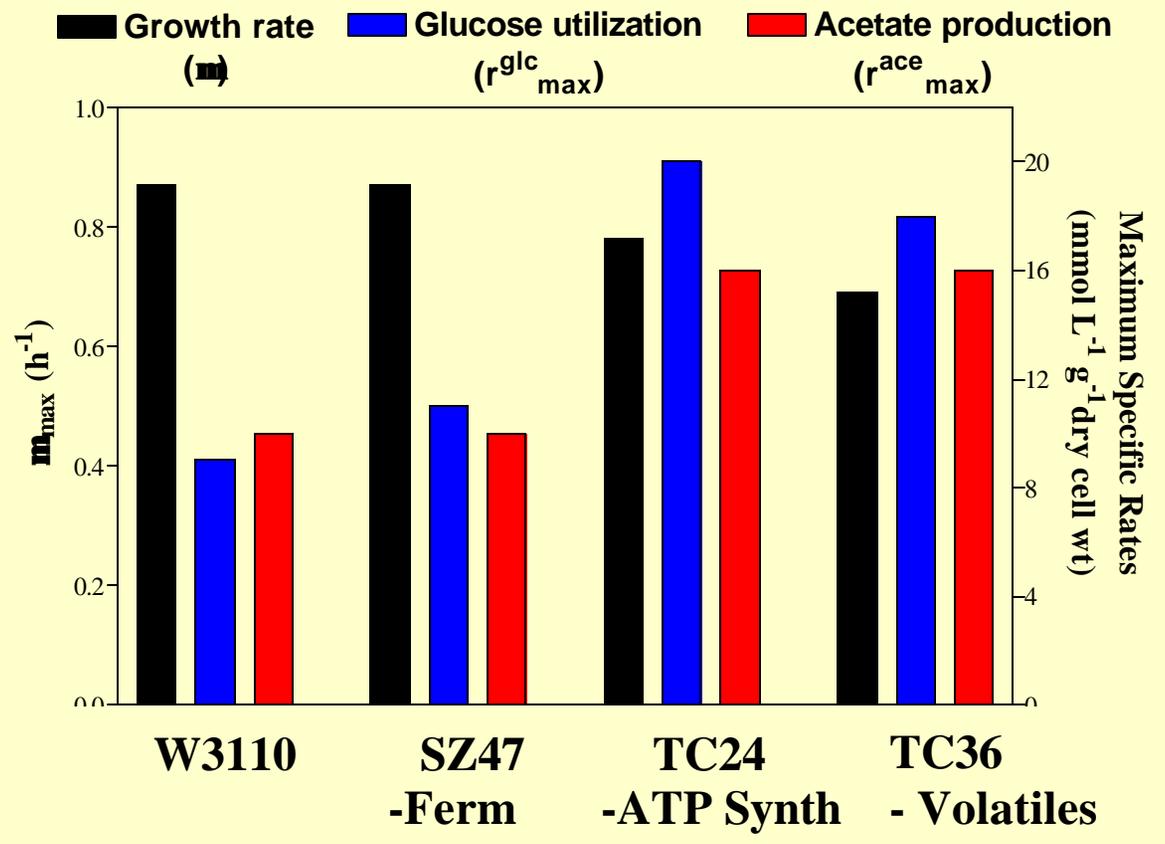
Strain SZ47 (*DfocA-pflB DfrdBC DldhA*) ~30 ATP/glu

Strain TC24(*DfocA-pflB DfrdBC DldhA DatpFH*) ~3ATP

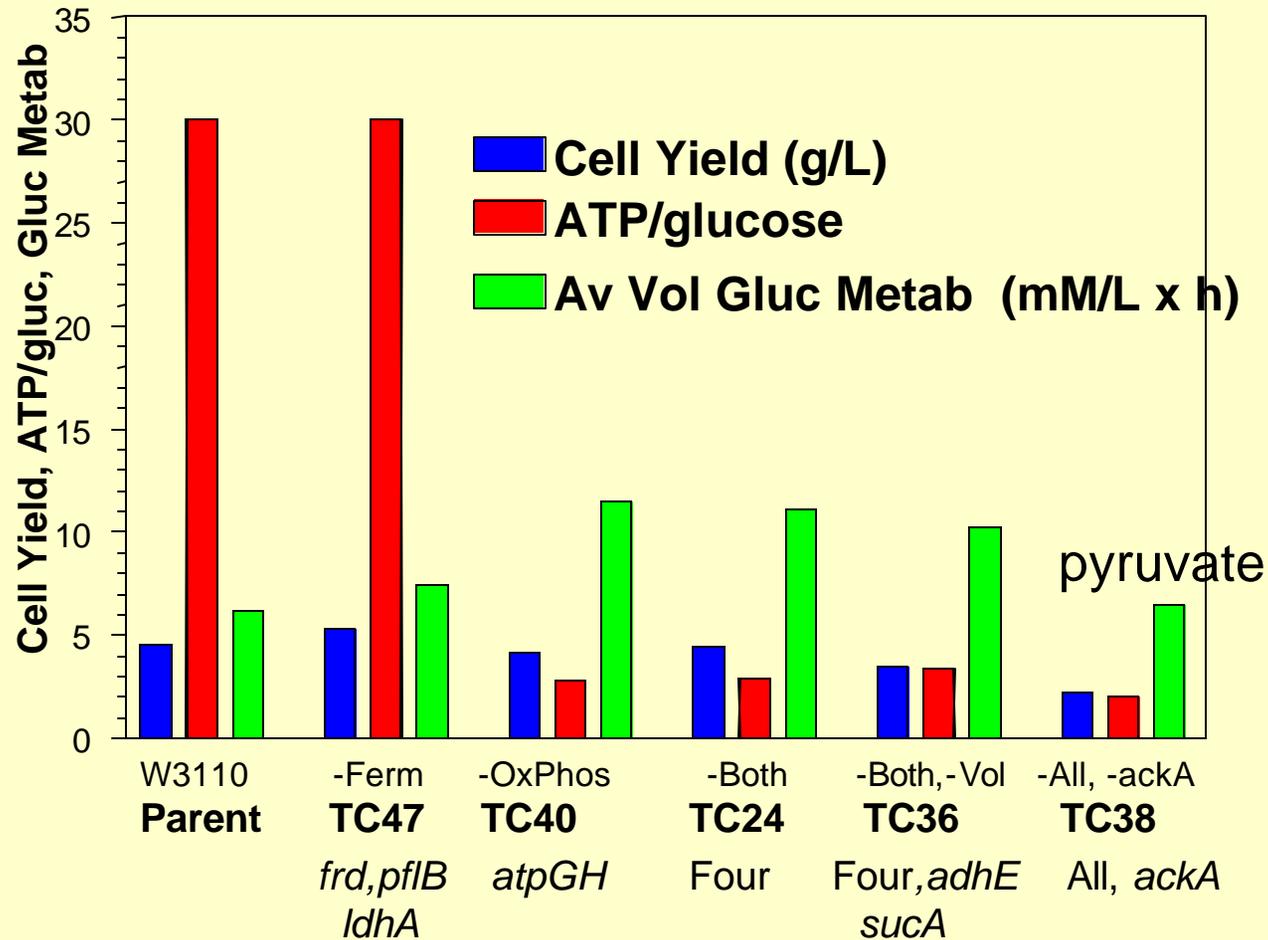
Strain TC36 (*DfocA-pflB DfrdBC DldhA DatpFH DadhE DsucA*) ~3.8 ATP/glucose



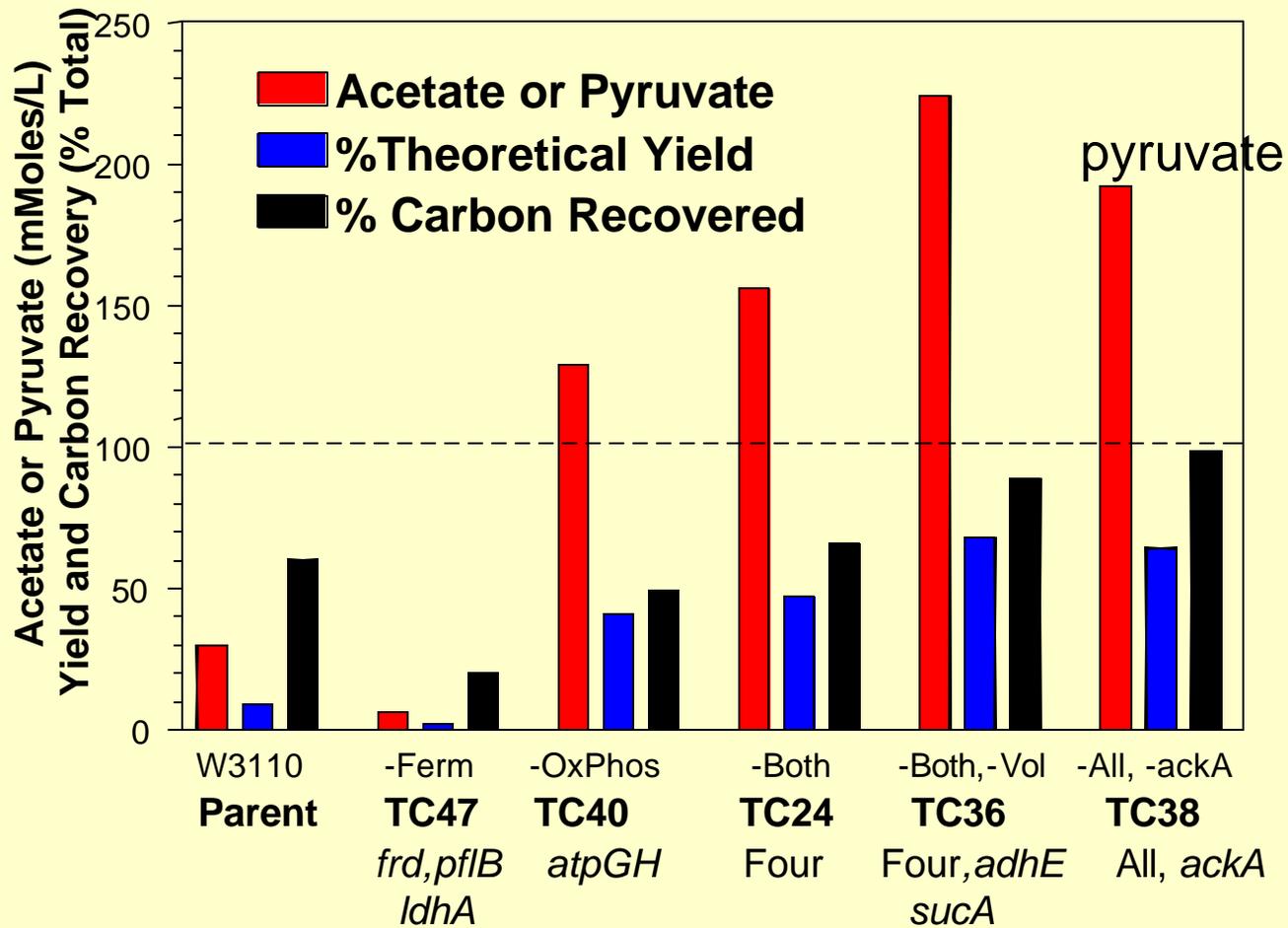
## Comparison of Maximum Specific Rates

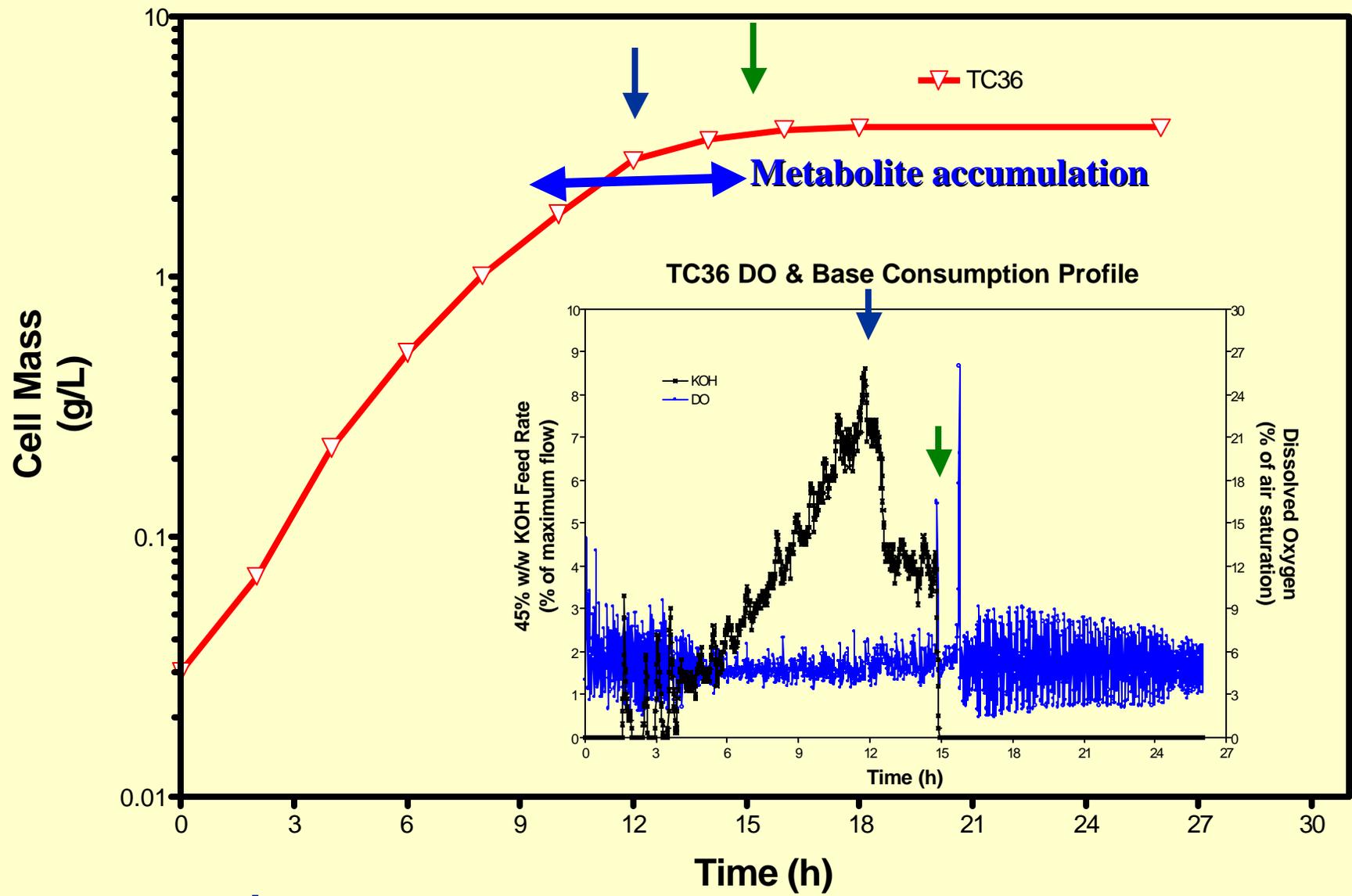


# Comparison of Strains

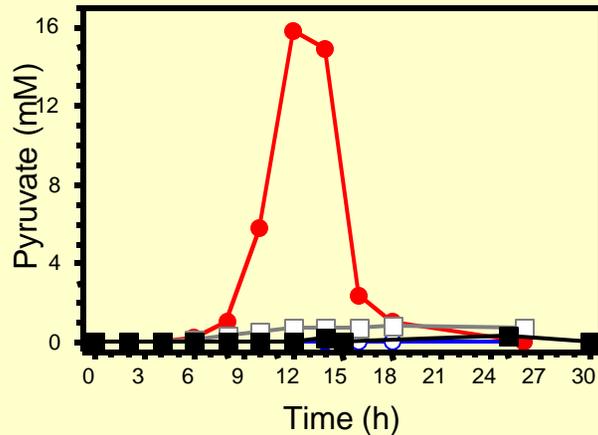


# Comparison of Strains

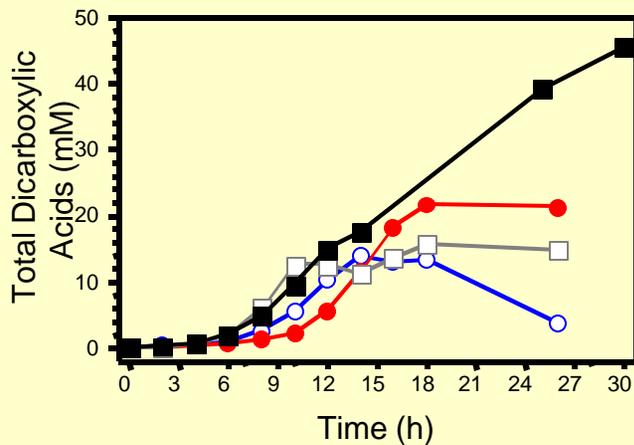




- ↓ Entrance into stationary phase
- ↓ Exhaustion of glucose

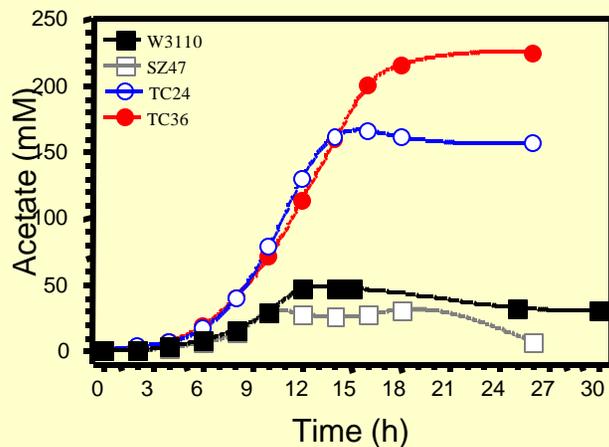


→ Accumulation of pyruvate indicates that glucose uptake & glycolysis may not be limiting acetate production.



→ W3110 accumulated ~ 3 fold higher concentrations of dicarboxylic acids than the engineered strains.

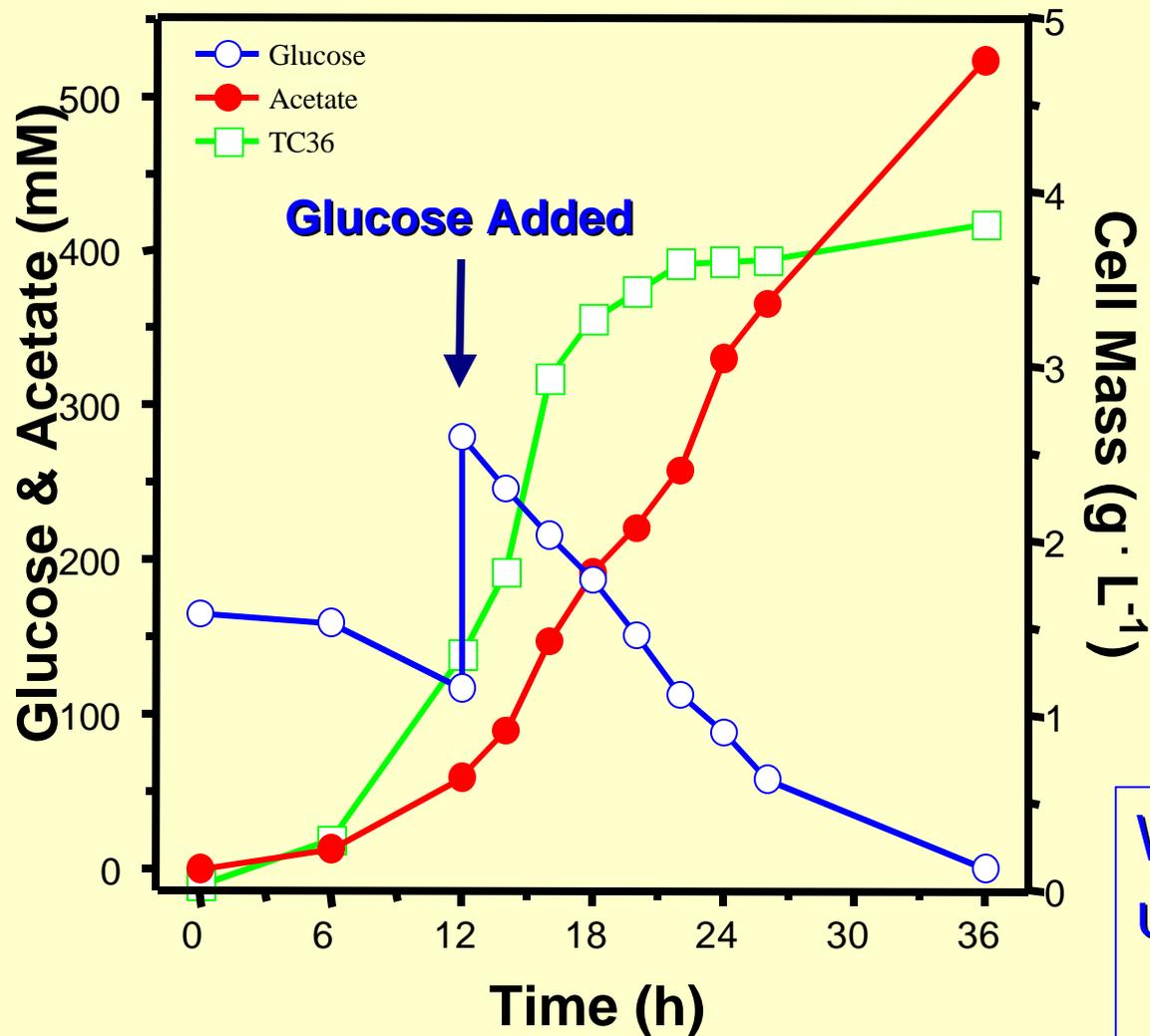
→ In general, accumulation of dicarboxylic acids was correlated with entry into stationary phase.



→ *DatpFH* resulted in ~25 fold increase in final acetate concentration/yield.

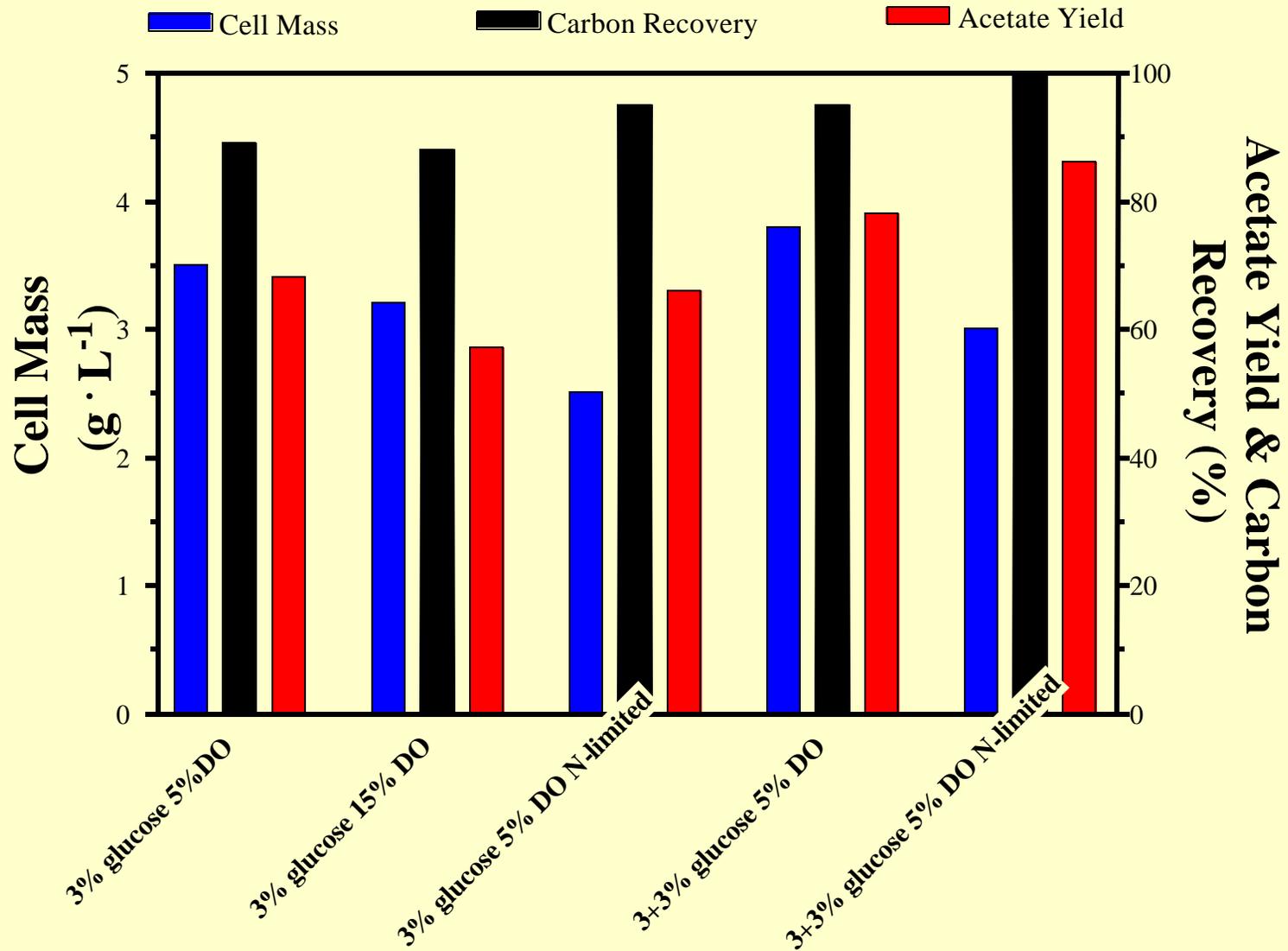
→ *DadhE DsucA* resulted in a further 1.4 fold increase in acetate concentration.

# Fermentation of 3% + 3% Glucose to Acetate



With glucose excess, up to 900 mM acetate was produced.

# Acetate Production Can Be Improved By Altering The Process Conditions (TC36)



# Conclusions

- *E. coli* can be engineered for efficient production of redox neutral & oxidized products (ace, pyr).
- $\Delta atpFH$  increased glycolytic flux by over 50%, acetate concentration and yield by 3-fold.
- $\Delta adhE \Delta sucA$  resulted in an additional increase in acetate concentration and yield, and improved carbon balance.
- The increase in glycolytic flux observed for TC24 & TC36 was attributed to the  $\Delta atpFH$  which reduced ATP production and provided gratuitous hydrolysis of excess ATP.
- Max acetate yields of 86% of theoretical; 10% of substrate carbon converted into biocatalyst
- Product stream relatively pure after cell removal